

FIG. 9

Seq ID NO: 11
MAGE-3 III CCTCCCAAGTCTCAGGAGCCCTCAGCCTCCCACTACCATGAACTACCTCTCTGAGCCAACTCATGAGGACTCCAGCAACCAAGAAAGAGG
MAGE-2 II ^{5601D NO: 9} CCTCCCAAGTCTCAGGAGCCCTCAGCCTCTCTGACTACCATCACTACTCTTGGAGACAACTCCGATGAGGGCTCCAGCAACCAAGAAAGAGG
MAGE-1 I ^{5601D NO: 8} CCTCCCAAGTCTCAGGAGCCCTCCGCTTTCCCACTACCATCACTCACTCACTGACAGAGGCAACCACTGAGGGTCCAGCAAGCCGTGAAAGAGG
225 CHO-8
III GGCCAAGCACTTCCCTGACC-TGGAGTCCGAGTTCCAAAGCACTCACTAGTAAAGGTGGCCGAGTTGGTTCAATTTCTGCTCCCAAGTATCGAGCCA
II GGCCAAGAACTGTTCCCGACCTTGGAGTCCGAGTTCCAAAGCACTCACTAGTAAAGGTGGCCGAGTTGGTTCAATTTCTGCTCCCAAGTATCGAGCCA
I GGCCAAGCACTTGGTATCC-TGGAGTCCGAGTTCCAAAGCACTCACTAGTAAAGGTGGCCGAGTTGGTTCAATTTCTGCTCCCAAGTATCGAGCCA
325
III GGGAGCCGCTCACAAAGGCAAAATGCTGGGAGTGTCTGAAATTTGCAATCTTCTCTGATCTTCAAGCAAGCTTCCAGTTCTTGCAGCT
II GGGAGCCGCTCACAAAGGCAAAATGCTGGGAGTGTCTGAAATTTGCAATCTTCTCTGATCTTCAAGCAAGCTTCCAGTTCTTGCAGCT
I GGGAGCACTCAAAAGGCAAAATGCTGGGAGTGTCTGAAATTTGCAATCTTCTCTGATCTTCCGAAAGCCCTGAGTCTTGCAGCT
425
III GGTCTTTGGCATCGAGTGTATGGAAGTGAACCCCATCGGCCACTGTAATCTTGGCACTGGCTTGGGCTCTCTACGATGGCTGCTGGGTGAAT
II GGTCTTTGGCATCGAGTGTGGAAGTGTATCCCATCAGCCACTGTACATCTTGTACCTGGCTTGGCTCTCTACGATGGCTGCTGGGTGAAT
I GGTCTTTGGCATCGAGTGTGGAAGTGTATCCCATCAGCCACTGTACATCTTGTACCTGGCTTGGCTCTCTACGATGGCTGCTGGGTGAAT
525
III CAGATCATGCCCAAGGAGGCTCTCTGATTAATCTCTGGCATATCGCAAGAGAGGGGCACTGTCCCTGAGGAGAAATCTGGGAGAGCTGAGTG
II CAGTCTATGCCCAAGAGAGGCTCTCTGATTAATCTCTGGCATATCGCAAGAGAGGGGCACTGTCCCTGAGGAGAAATCTGGGAGAGCTGAGTG
I CAGATCATGCCCAAGAGAGGCTCTCTGATTAATCTCTGGCATATCGCAAGAGAGGGGCACTGTCTCTGAGGAGAAATCTGGGAGAGCTGAGTG
625 CHO-9

